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L61 ANSWER 1 OF 7 HCAPLUS COPYRIGHT 2010 ACS on STN DUPLICATE 1
ACCESSION NUMBER:
                        2009:316615 HCAPLUS Full-text
DOCUMENT NUMBER:
                        151:418689
TITLE:
                        Strain typing of Mycoplasma cynos isolates from
                         dogs with respiratory disease
                         Mannering, Sally A.; McAuliffe, Laura; Lawes,
AUTHOR(S):
                         Joanna R.; Erles, Kerstin;
                         Brownlie, Joe
CORPORATE SOURCE:
                         The Royal Veterinary College, Hatfield, AL9 7TA,
SOURCE:
                        Veterinary Microbiology (2009), 135(3-4), 292-296
                        CODEN: VMICDQ; ISSN: 0378-1135
PUBLISHER:
                        Elsevier B.V.
DOCUMENT TYPE:
                        Journal
LANGUAGE:
                        English
     The association of Mycoplasma cynos with canine infectious respiratory disease
AB
     is increasingly being recognized. This study describes the strain typing of
     14 M. cynos isolates cultured from trachea and bronchoalveolar lavage samples
     of six dogs with respiratory disease, from two sep. kennels in the United
     Kingdom. The genetic similarity of the isolates was investigated using
     pulsed-field gel electrophoresis (PFGE) and random amplified polymorphic DNA
     (RAPD). Most of the isolates from four dogs housed at a re-homing kennel were
     genetically similar and some isolates from different dogs were
     indistinguishable by both PFGE and RAPD. These isolates were cultured from
     dogs with non-overlapping stays in the kennel, which may indicate maintenance
     of some strains within kennels. A small number of isolates showed much
     greater genetic heterogeneity and were genetically distinct from the main
     group of M. cynos strains. There was also a high degree of similarity of the
     M. cynos type strain (isolated from a dog with respiratory disease in Denmark
     in 1971) to at least one of the United Kingdom isolates using PFGE anal.,
     which may suggest possible conservation of pathogenic strains of M. cynos.
REFERENCE COUNT:
                         25
                               THERE ARE 25 CITED REFERENCES AVAILABLE FOR
                               THIS RECORD. ALL CITATIONS AVAILABLE IN THE
                               RE FORMAT
L61 ANSWER 2 OF 7 EMBASE COPYRIGHT (c) 2010 Elsevier B.V. All rights
     reserved on STN
                                                        DUPLICATE 2
ACCESSION NUMBER:
                    2007076462 EMBASE
                                          Full-text
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Serological evidence of Mycoplasma cynos infection in

TITLE:

canine infectious respiratory disease.

AUTHOR: Rycroft, Andrew N. (correspondence); Tsounakou,

Elizabeth; Chalker, Victoria

CORPORATE SOURCE: Department of Pathology and Infectious Diseases, Royal

Veterinary College, Hawkshead Lane, North Mymms, Herts

AL9 7TA, United Kingdom. arycroft@rvc.ac.uk

SOURCE: Veterinary Microbiology, (10 Mar 2007) Vol. 120, No.

3-4, pp. 358-362.

Refs: 24

ISSN: 0378-1135 CODEN: VMICDQ

PUBLISHER IDENT.: S 0378-1135(06)00458-5

COUNTRY: Netherlands
DOCUMENT TYPE: Journal; Article

FILE SEGMENT: 015 Chest Diseases, Thoracic Surgery and

Tuberculosis

004 Microbiology: Bacteriology, Mycology,

Parasitology and Virology

005 General Pathology and Pathological Anatomy

LANGUAGE: English SUMMARY LANGUAGE: English

ENTRY DATE: Entered STN: 13 Mar 2007

Last Updated on STN: 13 Mar 2007

Ab A high proportion of dogs suffer from respiratory disease when they are placed in kennels for vacation or re-homing. The role of Mycoplasma cynos as an initiating agent in canine infectious respiratory disease was investigated by examining the serological response of dogs to this organism at the time of entry into a large re-homing kennel. Forty-two paired serum samples from dogs (21-day interval) were examined for antibody to M. cynos using Western blotting. The development of antibody in the serum was related to clinical disease recorded over the same period. Sixty seven per cent of the dogs showed a two-fold or greater rise in antibody to M. cynos during the first 3 weeks in the kennel. Reactivity with a 45 kDa antigen was dominant. Of those showing a positive serological reaction, 80% had recorded clinical respiratory disease while 20% remained healthy. The findings of this study show that an antibody response to M. cynos is common in dogs entering the re-homing kennel and is positively related to the development of clinical respiratory disease. .COPYRGT. 2006 Elsevier B.V. All rights reserved.

L61 ANSWER 3 OF 7 HCAPLUS COPYRIGHT 2010 ACS on STN DUPLICATE 3

ACCESSION NUMBER: 2005:29227 HCAPLUS <u>Full-text</u>

DOCUMENT NUMBER: 142:133045

TITLE: Vaccines comprising attenuated viruses and bacteria or antigen-encoding nucleic acids and

antibodies for treating canine infectious

respiratory disease

INVENTOR(S): Brownlie, John; Chalker, Victoria

Jane; Erles, Kerstin

PATENT ASSIGNEE(S): The Royal Veterinary College, UK

SOURCE: PCT Int. Appl., 102 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent LANGUAGE: English

FAMILY ACC. NUM. COUNT: 1

PATENT INFORMATION:

PATENT NO. KIND DATE APPLICATION NO. DATE

WO 2005002618 A1 20050113 WO 2004-GB2865 20040701

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GB 2003-15323 A 20030701
                                              US 2007-849931
PRIORITY APPLN. INFO.:
                                               EP 2004-743211 A3 20040701
                                               WO 2004-GB2865 W 20040701
                                               US 2006-563199
                                                                     A3 20060901
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ASSIGNMENT HISTORY FOR US PATENT AVAILABLE IN LSUS DISPLAY FORMAT

AB A vaccine composition for vaccinating dogs comprising any one or more of (a) an agent capable of raising an immune response against Streptococcus equi sub species zooepidemicus in a dog, (b) an agent capable of raising an immune response against Mycoplasma cynos in a dog, and (c) an agent capable of raising an immune response against a Chlamydophila in a dog.

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OS.CITING REF COUNT: 1 THERE ARE 1 CAPLUS RECORDS THAT CITE THIS RECORD (1 CITINGS)
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REFERENCE COUNT: 16 THERE ARE 16 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

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L61 ANSWER 4 OF 7 HCAPLUS COPYRIGHT 2010 ACS on STN DUPLICATE 4 ACCESSION NUMBER: 2004:101304 HCAPLUS Full-text DOCUMENT NUMBER: 140:162357
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TITLE: Canine respiratory coronavirus (CRCV) spike protein, polymerase and hemagglutinin/esterase

gene and use thereof in diagnosis of and vaccine preparation against canine infectious respiratory

disease

INVENTOR(S): Brownlie, John; Chalker, Victoria

Jane; Erles, Kerstin

PATENT ASSIGNEE(S): The Royal Veterinary College, UK

SOURCE: PCT Int. Appl., 150 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent LANGUAGE: English

FAMILY ACC. NUM. COUNT: 1

PATENT INFORMATION:

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		2005533513						20051110 JP 2004-523905								0030701		
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ASSIGNMENT HISTORY FOR US PATENT AVAILABLE IN LSUS DISPLAY FORMAT

AB The present invention relates to a novel coronavirus, canine respiratory coronavirus (CRCV) identified from the respiratory tract kennelled dogs with canine infectious respiratory disease (CIRD). CRCV has a low level of homol. to the enteric canine coronavirus, but which has a high level of homol. to all bovine coronavirus strains (eq. Quebec and LY138) and human coronavirus strain OC43. Also

provided are amino acids specific to the CRCV polymerase, S protein and HE that are not present in the BCV, HCV and HEV S proteins. Serol. anal. also show that the presence of antibodies against CRCV on the day of entry into the kennel decreases the risk of developing respiratory disease. The CRCV spike, polymerase and hemagglutinin/esterase cDNA and protein partial sequences are listed in Figures (1) to (4), (13) and (14). CRCV might be useful to prepare vaccines against CIRD caused by CRCV as well as other causative agents.

OS.CITING REF COUNT: 3 THERE ARE 3 CAPLUS RECORDS THAT CITE THIS

RECORD (3 CITINGS)

REFERENCE COUNT: 10 THERE ARE 10 CITED REFERENCES AVAILABLE FOR

THIS RECORD. ALL CITATIONS AVAILABLE IN THE

RE FORMAT

L61 ANSWER 5 OF 7 HCAPLUS COPYRIGHT 2010 ACS on STN DUPLICATE 5

ACCESSION NUMBER: 2004:908070 HCAPLUS Full-text

DOCUMENT NUMBER: 142:130661

TITLE: Mycoplasmas associated with canine infectious

respiratory disease

AUTHOR(S): Chalker, Victoria J.; Owen, Wanda M. A.;

Paterson, Caren; Barker, Emily; Brooks, Harriet;

Rycroft, Andrew N.; Brownlie, Joe

CORPORATE SOURCE: Department of Pathology and Infectious Diseases,

Royal Veterinary College (RVC), University of

London, North Mymms, AL9 7TA, UK

SOURCE: Microbiology (Reading, United Kingdom) (2004),

150(10), 3491-3497

CODEN: MROBEO; ISSN: 1350-0872 Society for General Microbiology

DOCUMENT TYPE: Journal LANGUAGE: English

PUBLISHER:

AB Canine infectious respiratory disease (CIRD) is a complex infection that occurs worldwide predominantly in kennelled dogs, and several bacterial and viral microorganisms have been associated with outbreaks of CIRD. However, few studies have comprehensively examined the species of mycoplasma present in healthy dogs and those with CIRD. As part of an extensive study investigating the microorganisms involved in CIRD, the species of mycoplasma present throughout the respiratory tract of dogs with and without CIRD were determined Mycoplasmas were cultured from tonsillar, tracheal and bronchial lavage samples, and identified to the species level by PCR and sequencing. Mycoplasma cynos was demonstrated on the ciliated tracheal epithelium by in situ hybridization and was the only mollicute found to be associated with CIRD, but only in the lower respiratory tract. Isolation of M. cynos was correlated with an increased severity of CIRD, younger age and a longer time in the kennel.

OS.CITING REF COUNT: 3 THERE ARE 3 CAPLUS RECORDS THAT CITE THIS

RECORD (3 CITINGS)

REFERENCE COUNT: 18 THERE ARE 18 CITED REFERENCES AVAILABLE FOR

THIS RECORD. ALL CITATIONS AVAILABLE IN THE

RE FORMAT

L61 ANSWER 6 OF 7 MEDLINE on STN DUPLICATE 6

ACCESSION NUMBER: 2004292256 MEDLINE Full-text

DOCUMENT NUMBER: PubMed ID: 15193070

TITLE: Prevalence of Mycoplasma agassizii and Chelonian

herpesvirus in captive tortoises (Testudo sp.) in the

United Kingdom.

AUTHOR: Soares Jorge F; Chalker Victoria J;

Erles Kerstin; Holtby Sonya; Waters Michael;

McArthur Stuart

CORPORATE SOURCE: Department of Pathology and Infectious Diseases, Royal

Veterinary College, Hawkshead Lane, North Mymms,

Hertfordshire AL9 7TA, United Kingdom.

SOURCE: Journal of zoo and wildlife medicine: official

publication of the American Association of Zoo

Veterinarians, (2004 Mar) Vol. 35, No. 1, pp. 25-33.

Journal code: 8915208. ISSN: 1042-7260. L-ISSN:

1042-7260.

PUB. COUNTRY: United States

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

(RESEARCH SUPPORT, NON-U.S. GOV'T)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 200411

ENTRY DATE: Entered STN: 15 Jun 2004

Last Updated on STN: 6 Nov 2004 Entered Medline: 6 Nov 2004

AΒ During the months of April to August in 1999 and 2002, oral swabs were collected from 146 tortoises (Testudo sp.) in private collections in the United Kingdom and tested by polymerase chain reaction (PCR) for the presence of Mycoplasma agassizii and Chelonian herpesvirus (ChHV). The presence of M. agassizii was confirmed by restriction digestion of the PCR product. A 307-bp fragment of the ChHV UL5 homologue gene was sequenced and found to show most similarity to equine herpesvirus type 1. A prevalence of 15.8 and 8.2% was found for M. agassizii and ChHV, respectively. Comparison of the carriage of both M. agassizii and ChHV in different species of tortoises correlated the presence of M. agassizii with Testudo horsfieldii and ChHV with Testudo marginata and Testudo graeca iberia. An association of ChHV with stomatitis was also found. Mixed infections with both agents were detected. The findings further demonstrate this pathogen-tortoise association and the cross transmission of these infections if different tortoise species are housed together.

L61 ANSWER 7 OF 7 BIOSIS COPYRIGHT (c) 2010 The Thomson Corporation on

STN

ACCESSION NUMBER: 2002:212345 BIOSIS Full-text

DOCUMENT NUMBER: PREV200200212345

TITLE: The role of Mycoplasmas in infectious canine

tracheobronchitis.

AUTHOR(S): Chalker, V. J. [Reprint author]; Toomey, C.

[Reprint author]; %xles, %. [Reprint author]; Brooks, H. [Reprint author]; Opperman, S. [Reprint author]; Rycroft, A. [Reprint author]; %xownlie,

J. [Reprint author]

CORPORATE SOURCE: Royal Veterinary College, Hatfield, Hertfordshire, UK

SOURCE:

Abstracts of the General Meeting of the American Society for Microbiology, (2001) Vol. 101, pp. 392.

print.

Meeting Info.: 101st General Meeting of the American Society for Microbiology. Orlando, FL, USA. May 20-24,

2001. American Society of Microbiology.

ISSN: 1060-2011.

DOCUMENT TYPE: Conference; (Meeting)

Conference; Abstract; (Meeting Abstract)

LANGUAGE: English

ENTRY DATE: Entered STN: 27 Mar 2002

Last Updated on STN: 27 Mar 2002

AB Infectious canine trachiobronchitis (ICT, kennel cough) is a major problem in kennelled dogs world-wide, particularly when dogs are housed in a high density environment. The disease process begins with mild nasal discharge and coughing, but can develop and lead to severe depression, secondary infection,

and in some cases even death. The causative agents of ICT have never been accurately determined and infection is thought to occur as a complex of several infectious agents, including Bordetella bronchiseptica and Canine Parinfluenza Virus. The precise role of Mycoplasma species in the aetiology of canine respiratory disease has never been characterised. In collaboration with a well-established kennel (apprx3000 animals) with a history of severe and recurrent ICT, we sought to evaluate which Mycoplasma species were present in both the upper and lower canine respiratory tract. Analysis of clinically normal dogs and those in all stages of respiratory disease led to the discovery that a large number of Mycoplasma species are present in the canine respiratory tract, and that total Mycoplasma load increases with increasing disease severity. Isolates were initially grouped with RAPD analysis and then further characterised by sequencing of the 16S rRNA gene concurrent with growth inhibition testing. In addition, we correlate the presence of Mycoplasmas to ICT disease severity and clinical manifestations.

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